IAP20 Rec'd PCT/PTO 22 JUN 2006

1

SEQUENCE LISTING

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<110> KIM, JIN-SOO
      PARK, KYUNG-SOON
     JANG, YOUNG-SOON
<120> REGULATION OF PROKARYOTIC GENE EXPRESSION WITH ZINC
      FINGER PROTEINS
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<150> PCT/KR04/003420
<151> 2004-12-23
<150> 60/532,362
<151> 2003-12-23
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Thr Gln His Arg Arg Ile His 20

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Val Cys Gly Lys Ala Phe Arg His Ser Ser Ser Leu Val Arg His Gln 35 40 45

Arg Thr His Thr Gly Glu Lys Pro Tyr Arg Cys Lys Tyr Cys Asp Arg 50 55 60

Ser Phe Ser Ile Ser Ser Asn Leu Gln Arg His Val Arg Asn Ile His 65 70 75 80

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Lys Gln His Thr Arg Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Lys 20 25 30

Gln Cys Gly Lys Ala Phe Gly Cys Pro Ser Asn Leu Arg Arg His Gly 35 40 45

Arg Thr His Thr Gly Glu Lys Pro Tyr Arg Cys Lys Tyr Cys Asp Arg 50 55 60

Ser Phe Ser Ile Ser Ser Asn Leu Gln Arg His Val Arg Asn Ile His 65 70 75 80

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<212> PRT

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<223> Description of Artificial Sequence: Synthetic protein

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1 5 10 15

Gln Arg His Val Arg Asn Ile His Thr Gly Glu Lys Pro Phe Gln Cys
20 25 30

Lys Thr Cys Gln Arg Lys Phe Ser Arg Ser Asp His Leu Lys Thr His 35 40 45

Thr Arg Thr His Thr Gly Glu Lys Pro Tyr Glu Cys His Asp Cys Gly 50 60

Lys Ser Phe Arg Gln Ser Thr His Leu Thr Arg His Arg Arg Ile His 65 70 75 80

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Val Ser Ser Thr Leu Ile Arg His Gln Arg Ile His 100 105

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Asn Val His Lys Arg Thr His Thr Gly Glu Lys Pro Tyr Glu Cys Asp 20 25 30 His Cys Gly Lys Ala Phe Ser Val Ser Ser Asn Leu Asn Val His Arg 35 40 45

Arg Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Glu Glu Cys Gly Lys
50 55 60

Ala Phe Thr Gln Ser Ser Asn Leu Thr Lys His Lys Lys Ile His Thr 65 70 75 80

Gly Glu Lys Pro Tyr Lys Cys Glu Glu Cys Gly Lys Ala Phe Thr Gln 85 90 95

Ser Ser Asn Leu Thr Lys His Lys Lys Ile His
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His Cys Gly Lys Ser Phe Ser Gln Ser Ser His Leu Asn Val His Lys 35 40 45

Arg Thr His Thr Gly Ġlu Lys Pro Tyr Glu Cys His Asp Cys Gly Lys 50 60

Ser Phe Arg Gln Ser Thr His Leu Thr Arg His Arg Arg Ile His Thr 65 70 75 80

Gly Glu Lys Pro Tyr Lys Cys Pro Asp Cys Gly Lys Ser Phe Ser Gln 85 90 95

Ser Ser Ser Leu Ile Arg His Gln Arg Thr His 100 105

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His Cys Gly Lys Ser Phe Ser Gln Ser Ser His Leu Asn Val His Lys 35 40 45

Arg Thr His Thr Gly Glu Lys Pro Tyr Glu Cys Asp His Cys Gly Lys
50 55 60

Ala Phe Ser Val Ser Ser Asn Leu Asn Val His Arg Arg Ile His Thr
65 70 75 80

Gly Glu Lys Pro Tyr Glu Cys Asp His Cys Gly Lys Ser Phe Ser Gln 85 90 95

Ser Ser His Leu Asn Val His Lys Arg Thr His 100 105

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Thr Arg His Arg Arg Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Met 20 25 30

Glu Cys Gly Lys Ala Phe Asn Arg Arg Ser His Leu Thr Arg His Gln 35 40 45

Arg Ile His Thr Gly Glu Lys Pro Tyr Glu Cys His Asp Cys Gly Lys
50 55 60

Ser Phe Arg Gln Ser Thr His Leu Thr Arg His Arg Arg Ile His Thr 65 70 75 80

Gly Glu Lys Pro Tyr Glu Cys His Asp Cys Gly Lys Ser Phe Arg Gln 85 90 95

Ser Thr His Leu Thr Arg His Arg Arg Ile His
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Cys Lys Thr Cys Gln Arg Lys Phe Ser Arg Ser Asp His Leu Lys Thr 35 40 45

His Thr Arg Thr His Thr Gly Glu Lys Pro Tyr Glu Cys Asp His Cys 50 55 60

Gly Lys Ser Phe Ser Gln Ser Ser His Leu Asn Val His Lys Arg Thr 65 70 75 80

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Thr Gln His Arg Arg Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Met 20 25 30

Glu Cys Gly Lys Ala Phe Asn Arg Arg Ser His Leu Thr Arg His Gln 35 40 45

Arg Ile His Thr Gly Glu Lys Pro Tyr Glu Cys His Asp Cys Gly Lys
50 55 60

Ser Phe Arg Gln Ser Thr His Leu Thr Arg His Arg Arg Ile His Thr 65 70 75 80

Gly Glu Lys Pro Tyr Glu Cys His Asp Cys Gly Lys Ser Phe Arg Gln \$85\$

Ser Thr His Leu Thr Arg His Arg Arg Ile His 100 105

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His Cys Gly Lys Ala Phe Ser Val Ser Ser Asn Leu Asn Val His Arg

Arg Ile His Thr Gly Glu Lys Pro Phe Glu Cys Lys Asp Cys Gly Lys
50 55 60

Ala Phe Ile Gln Lys Ser Asn Leu Ile Arg His Gln Arg Thr His Thr 65 70 75 80

Gly Glu Lys Pro Tyr Lys Cys Lys Gln Cys Gly Lys Ala Phe Gly Cys 85 90 95

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Asp Cys Gly Lys Ser Phe Arg Gln Ser Thr His Leu Thr Arg His Arg 35 40 45

Arg Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Pro Asp Cys Gly Lys 50 55 60

Ser Phe Ser Gln Ser Ser Ser Leu Ile Arg His Gln Arg Thr His Thr 65 70 75 80

Gly Glu Lys Pro Phe Gln Cys Lys Thr Cys Gln Arg Lys Phe Ser Arg 85 90 95

Ser Asp His Leu Lys Thr His Thr Arg Thr His
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His Cys Gly Lys Ser Phe Ser Gln Ser Ser His Leu Asn Val His Lys 35 40 45

Arg Thr His Thr Gly Glu Lys Pro Tyr Glu Cys His Asp Cys Gly Lys 50 55 60

Ser Phe Arg Gln Ser Thr His Leu Thr Arg His Arg Arg Ile His Thr 65 70 75 80

Gly Glu Lys Pro Phe Glu Cys Lys Asp Cys Gly Lys Ala Phe Ile Gln 85 90 95

Lys Ser Asn Leu Ile Arg His Gln Arg Thr His \$100\$

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<223> Description of Artificial Sequence: Synthetic protein

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Ala Leu Ala Arg His Lys Arg Thr His Thr Gly Glu Lys Pro Phe Gln 20 25 30

Cys Lys Thr Cys Gln Arg Lys Phe Ser Arg Ser Asp His Leu Lys Thr $35 \hspace{1cm} 40 \hspace{1cm} 45$

His Thr Arg Thr His Thr Gly Glu Lys Pro Tyr Lys Cys Glu Glu Cys
50 60

Gly Lys Ala Phe Thr Gln Ser Ser Asn Leu Thr Lys His Lys Lys Ile 65 70 75 80

His Thr Gly Glu Lys Pro Tyr Glu Cys His Asp Cys Gly Lys Ser Phe 85 90 95

Arg Gln Ser Thr His Leu Thr Arg His Arg Arg Ile His

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<213> Simian parainfluenza virus 5

<400> 57

Gly Lys Pro Ile Pro Asn Pro Leu Leu Gly Leu Asp Ser

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Arg Ser Asp Glu Leu Thr Arg His Ile Arg Ile His Thr Gly Gln Lys 20 25 30

Pro Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser Arg Ser Asp His 35 40 45

Leu Thr Thr His Ile Arg Thr His Thr Gly Glu Lys Pro Phe Ala Cys 50 55 60

Asp Ile Cys Gly Arg Lys Phe Ala Arg Ser Asp Glu Arg Lys Arg His

Thr Lys Ile His Leu Arg Gln Lys Asp

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Arg Arg His Gly Arg Thr His 20

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His Arg His Gln Arg Thr His 20

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Arg Arg His Cys Ile Leu His 20

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Asn Arg His Arg Arg Thr His 20

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Val Arg His Gln Arg Thr His
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Gln Arg His Val Arg Asn Ile His
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Asn Val His Arg Arg Ile His
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Ile Arg His His Lys Leu His
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Arg Arg His Glu Lys Thr His
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Arg Arg His Glu Lys Thr His
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 Val Arg His Lys Arg Thr His
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Ile Ile His Gln Arg Thr His
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Thr Lys His Lys Lys Ile His
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<210> 81
<211> 23
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<213> Homo sapiens
Phe Glu Cys Lys Asp Cys Gly Lys Ala Phe Ile Gln Lys Ser Asn Leu
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Ile Arg His Gln Arg Thr His 20
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<210> 82

<211> 23

<212> PRT <213> Homo sapiens

<400> 82

Tyr Val Cys Arg Glu Cys Arg Arg Gly Phe Ser Gln Lys Ser Asn Leu 1 5 10 15

Ile Arg His Gln Arg Thr His 20

<210> 83

<211> 23

<212> PRT

<213> Homo sapiens

<400> 83

Tyr Glu Cys Glu Lys Cys Gly Lys Ala Phe Asn Gln Ser Ser Asn Leu 1 5 10 15

Thr Arg His Lys Lys Ser His

<210> 84

<211> 23

<212> PRT

<213> Homo sapiens

<400> 84

Tyr Glu Cys Asn Thr Cys Arg Lys Thr Phe Ser Gln Lys Ser Asn Leu 1 5 10 15

Ile Val His Gln Arg Thr His

<210> 85

<211> 23

<212> PRT

<213> Homo sapiens

<400> 85

Tyr Val Cys Ser Lys Cys Gly Lys Ala Phe Thr Gln Ser Ser Asn Leu 1 5 10

Thr Val His Gln Lys Ile His

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<210> 86
<211> 23
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<213> Homo sapiens
<400> 86
Tyr Lys Cys Asp Glu Cys Gly Lys Asn Phe Thr Gln Ser Ser Asn Leu
Ile Val His Lys Arg Ile His
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<210> 87
<211> 23
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Gly Val His Gln Arg Thr His
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<210> 88
<211> 23
<212> PRT
<213> Homo sapiens
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Tyr Glu Cys Val Gln Cys Gly Lys Gly Phe Thr Gln Ser Ser Asn Leu
Ile Thr His Gln Arg Val His
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<210> 89
<211> 23
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<213> Homo sapiens
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Tyr Lys Cys Pro Asp Cys Gly Lys Ser Phe Ser Gln Ser Ser Ser Leu
Ile Arg His Gln Arg Thr His
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<210> 90
<211> 23
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<213> Homo sapiens
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<213> Homo sapiens
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Ile Arg His Arg Arg Ser His
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<211> 23
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Thr Arg His Lys Ile Val His
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<210> 93
<211> 23
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<213> Homo sapiens
Tyr Glu Cys Asn Glu Cys Gly Lys Ala Phe Ala Gln Asn Ser Thr Leu
Arg Val His Gln Arg Ile His
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<210> 94
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<213> Homo sapiens
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Tyr Glu Cys His Asp Cys Gly Lys Ser Phe Arg Gln Ser Thr His Leu
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Thr Gln His Arg Arg Ile His
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<211> 23
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<213> Homo sapiens
<400> 95
Tyr Glu Cys His Asp Cys Gly Lys Ser Phe Arg Gln Ser Thr His Leu
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Thr Arg His Arg Arg Ile His
<210> 96
<211"> 22"
<212> PRT
<213> Homo sapiens
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His Lys Cys Leu Glu Cys Gly Lys Cys Phe Ser Gln Asn Thr His Leu
Thr Arg His Gln Arg Thr
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<210> 97
<211> 25
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<213> Homo sapiens
Tyr Val Cys Asp Val Glu Gly Cys Thr Trp Lys Phe Ala Arg Ser Asp
Glu Leu Asn Arg His Lys Lys Arg His
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<210> 98
<211> 25
<212> PRT
<213> Homo sapiens
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Tyr His Cys Asp Trp Asp Gly Cys Gly Trp Lys Phe Ala Arg Ser Asp
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Glu Leu Thr Arg His Tyr Arg Lys His
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<210> 99
<211> 25
<212> PRT
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<213> Homo sapiens

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<400> 99
Tyr Arg Cys Ser Trp Glu Gly Cys Glu Trp Arg Phe Ala Arg Ser Asp
Glu Leu Thr Arg His Phe Arg Lys His
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<210> 100
<211> 25
<212> PRT
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Glu Leu Ser Arg His Arg Arg Thr His
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<210> 101
<211> 25
<212> PRT
<213> Homo sapiens
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Phe Ala Cys Ser Trp Gln Asp Cys Asn Lys Lys Phe Ala Arg Ser Asp
                                     10
Glu Leu Ala Arg His Tyr Arg Thr His
             20
<210> 102
<211> 25
<212> PRT
<213> Homo sapiens
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Tyr His Cys Asn Trp Asp Gly Cys Gly Trp Lys Phe Ala Arg Ser Asp
Glu Leu Thr Arg His Tyr Arg Lys His
            20
<210> 103
<211> 24
<212> PRT
<213> Homo sapiens
Phe Leu Cys Gln Tyr Cys Ala Gln Arg Phe Gly Arg Lys Asp His Leu
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Thr Arg His Met Lys Lys Ser His 20

<210> 104

<211> 23

<212> PRT

<213> Homo sapiens

<400> 104

Phe Gln Cys Lys Thr Cys Gln Arg Lys Phe Ser Arg Ser Asp His Leu 1 5 10 15

Lys Thr His Thr Arg Thr His 20

<210> 105

<211> 23

<212> PRT

<213> Homo sapiens

<400> 105

Phe Ala Cys Glu Val Cys Gly Val Arg Phe Thr Arg Asn Asp Lys Leu
1 5 10 15

Lys Ile His Met Arg Lys. His

<210> 106

<211> 25

<212> PRT

<213> Homo sapiens

<400> 106

Tyr Val Cys Asp Val Glu Gly Cys Thr Trp Lys Phe Ala Arg Ser Asp 1 5 10 15

Lys Leu Asn Arg His Lys Lys Arg His 20 25

<210> 107

<211> 23

<212> PRT

<213> Homo sapiens

<400> 107

Tyr Lys Cys Met Glu Cys Gly Lys Ala Phe Asn Arg Arg Ser His Leu 1 5 10 15

Thr Arg His Gln Arg Ile His 20

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<210> 108
<211> 23
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<213> Homo sapiens
<400> 108
Tyr Ile Cys Arg Lys Cys Gly Arg Gly Phe Ser Arg Lys Ser Asn Leu
Ile Arg His Gln Arg Thr His
            20
<210> 109
<211> 23
<212> PRT
<213> Homo sapiens
                                     - --
                                                     3 - 3 - 3 -
<400> 109
Tyr Leu Cys Ser Glu Cys Asp Lys Cys Phe Ser Arg Ser Thr Asn Leu
                      . 10
1 5
Ile Arg His Arg Arg Thr His
<210> 110
<211> 23
<212> PRT
<213> Homo sapiens
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Tyr Glu Cys Lys Glu Cys Gly Lys Ala Phe Ser Ser Gly Ser Asn Phe
                                  10
Thr Arg His Gln Arg Ile His
           20
<210> 111
<211> 23
<212> PRT
<213> Homo sapiens
Tyr Glu Cys Asp His Cys Gly Lys Ala Phe Ser Val Ser Ser Asn Leu
Asn Val His Arg Arg Ile His
            20
<210> 112
<211> 23
<212> PRT
<213> Homo sapiens
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Tyr Thr Cys Lys Gln Cys Gly Lys Ala Phe Ser Val Ser Ser Ser Leu
                                    10
Arg Arg His Glu Thr Thr His
<210> 113
<211> 23
<212> PRT
<213> Homo sapiens
<400> 113
Tyr Glu Cys Asn Tyr Cys Gly Lys Thr Phe Ser Val Ser Ser Thr Leu
                                                     15
            5 10
Ile Arg His Gln Arg Ile His
            20
<210> 114
<211> 23
<212> PRT
<213> Homo sapiens
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Tyr Arg Cys Glu Glu Cys Gly Lys Ala Phe Arg Trp Pro Ser Asn Leu
                                    10
                 5
Thr Arg His Lys Arg Ile His
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<210> 115
<211> 6
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<222> (3)
<223> Glu or Gln
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<221> MOD_RES
<222> (4)
<223> Lys or Arg
<220>
<221> MOD_RES
<222> (6)
<223> Tyr or Phe
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Thr Gly Xaa Xaa Pro Xaa
<210> 116
<211> 28
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<223> Phe or Tyr
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<222> (2)
<223> variable amino acid
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<222> (4)..(8)
<223> region may encompass 2-5 variable amino acids
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<223> Phe or Tyr
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<222> (16)
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<222> (19)
<223> hydrophobic amino acid
<220>
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<222> (20)
<223> variable amino acid
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<222> (23)..(27)
<223> region may encompass 3-5 variable amino acids
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Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Cys Xaa
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Ser Asn Xaa Xaa Arg His Xaa Xaa Xaa Xaa His
             20
<210> 117
<211> 267
<212> DNA
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taagcctatc cctaaccctc tcctcggtct cgattctaca caagctatgg gtgctcctcc 180
aaaaaagaag agaaaggtag ctggatccac tagtaacggc cgccagtgtg ctggaattct 240
gcagatatcc atcacactgg cggccgc
<210> 118
<211> 25
<212> PRT
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Ala Leu Ala Arg His Lys Arg Thr His
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<210> 119
<211> 23
<212> PRT
<213> Homo sapiens
<400> 119
Tyr Lys Cys Lys Gln Cys Gly Lys Ala Phe Gly Cys Pro Ser Asn Leu
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Arg Arg His Gly Arg Thr His 20 <210> 120 <211> 23
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<212> PRT <213> Homo sapiens

<400> 120

Tyr Thr Cys Ser Asp Cys Gly Lys Ala Phe Arg Asp Lys Ser Cys Leu 1 5 10

Asn Arg His Arg Arg Thr His 20

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<223> Description of Artificial Sequence: Synthetic peptide

Asn Leu Thr Arg His Ile Arg Ile His
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<210> 122 <211> 23 <212> PRT <213> Homo sapiens

Val Arg His Gln Arg Thr His 20

<210> 123 <211> 24 <212> PRT <213> Homo sapiens

Gln Arg His Val Arg Asn Ile His 20

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<210> 124
<211> 23
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<213> Homo sapiens
Tyr Lys Cys His Gln Cys Gly Lys Ala Phe Ile Gln Ser Phe Asn Leu
1
Arg Arg His Glu Arg Thr His
<210> 125
<211>-23----
<212> PRT
<213> Drosophila sp.
<400> 125
Tyr Thr Cys Ser Tyr Cys Gly Lys Ser Phe Thr Gln Ser Asn Thr Leu
Lys Gln His Thr Arg Ile His
             20
<210> 126
<211> 23
<212> PRT
<213> Homo sapiens
<400> 126
Tyr Glu Cys Asp His Cys Gly Lys Ser Phe Ser Gln Ser Ser His Leu
Asn Val His Lys Arg Thr His
             20.
<210> 127
<211> 23
<212> PRT
<213> Homo sapiens
<400> 127
Tyr Met Cys Ser Glu Cys Gly Arg Gly Phe Ser Gln Lys Ser Asn Leu
                                      10
Ile Ile His Gln Arg Thr His
             20
<210> 128
<211> 23
<212> PRT
<213> Homo sapiens
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Tyr Lys Cys Glu Glu Cys Gly Lys Ala Phe Thr Gln Ser Ser Asn Leu
Thr Lys His Lys Lys Ile His
            20
<210> 129
<211> 23
<212> PRT
<213> Homo sapiens
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Phe Glu Cys Lys Asp Cys Gly Lys Ala Phe Ile Gln Lys Ser Asn Leu
Ile Arg His Gln Arg Thr His
            20
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 Tyr Val Cys Ser Lys Cys Gly Lys Ala Phe Thr Gln Ser Ser Asn Leu
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 Thr Val His Gln Lys Ile His
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 <213> Homo sapiens
 Tyr Lys Cys Pro Asp Cys Gly Lys Ser Phe Ser Gln Ser Ser Leu
 Ile Arg His Gln Arg Thr His
            20
 <210> 132
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 Tyr Glu Cys His Asp Cys Gly Lys Ser Phe Arg Gln Ser Thr His Leu
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38
Thr Gln His Arg Arg Ile His
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<210> 133
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<212> PRT
<213> Homo sapiens
<400> 133
Tyr Glu Cys His Asp Cys Gly Lys Ser Phe Arg Gln Ser Thr His Leu
Thr Arg His Arg Arg Ile His
                                                and the contract of the contra
<210> 134
<211> 23
<212> PRT
<213> Homo sapiens
<400> 134
Phe Gln Cys Lys Thr Cys Gln Arg Lys Phe Ser Arg Ser Asp His Leu
Lys Thr His Thr Arg Thr His
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 <211> 25
 <212> PRT
 <213> Homo sapiens
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 <212> PRT
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peptide

Phe Ala Cys Pro Glu Cys Pro Lys Arg Phe Met Arg Ser Asp Asn Leu

<223> Description of Artificial Sequence: Synthetic

Thr Gln His Ile Lys Thr His

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<213> Homo sapiens
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Thr Arg His Gln Arg Ile His
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<211> 23
<212> PRT
<213> Homo sapiens
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Ile Arg His Gln Arg Thr His
<210> 139
<211> 23
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Asn Val His Arg Arg Ile His
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<211> 23
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<213> Homo sapiens
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Tyr Thr Cys Lys Gln Cys Gly Lys Ala Phe Ser Val Ser Ser Leu
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Arg Arg His Glu Thr Thr His
             20
<210> 141
<211> 23
<212> PRT
<213> Homo sapiens
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Tyr Glu Cys Asn Tyr Cys Gly Lys Thr Phe Ser Val Ser Ser Thr Leu
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Ile Arg His Gln Arg Ile His
<210> 142
<211> 23
<212> PRT
<213> Homo sapiens
<400> 142
Tyr Arg Cys Glu Glu Cys Gly Lys Ala Phe Arg Trp Pro Ser Asn Leu
1.0
Thr Arg His Lys Arg Ile His
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<210> 143 .
<211> 12
<212> DNA
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<223> Description of Artificial Sequence: Synthetic
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daadaaaath ga
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<220>
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<400> 144
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gyagrahgan ggk
<210> 145
<211> 12
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<213> Artificial Sequence

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<400> 145
hgaaathgag gt
                                                                                                                                                                                                                                                         12
<210> 146
<211> 12
<212> DNA
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                 oligonucleotide
                                          والمتعال المرازي والمرازي والمرازي والمتعال في المتعال المتعال المتعال والمتعال والم
<400> 146
                                                                                                                                                                                                                                                          12
gragragggg ra
 <210> 147
 <211> 12
 <212> DNA
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 grahganggg tc
  <210> 148
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  gragragggh ga
  <210> 149
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 gavgaaaath ga
 <210> 150
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 <400> 150
                                                               12
 ngggyagraa at
 <210> 151
 <211> 13
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 gaagrahgan ggk
 <210> 152
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gradaanggg tc
<210> 153
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<400> 153
                                                                   12
gaagrahgan gg
<210> 154
<211> 189
<212> PRT
<213> Escherichia coli
<400> 154
Met Lys Arg Leu Ile Val Gly Ile Ser Gly Ala Ser Gly Ala Ile Tyr
Gly Val Arg Leu Leu Gln Val Leu Arg Asp Val Thr Asp Ile Glu Thr
His Leu Val Met Ser Gln Ala Ala Arg Gln Thr Leu Ser Leu Glu Thr
Asp Phe Ser Leu Arg Glu Val Gln Ala Leu Ala Asp Val Thr His Asp
Ala Arg Asp Ile Ala Ala Ser Ile Ser Ser Gly Ser Phe Gln Thr Leu
Gly Met Val Ile Leu Pro Cys Ser Ile Lys Thr Leu Ser Gly Ile Val
His Ser Tyr Thr Asp Gly Leu Leu Thr Arg Ala Ala Asp Val Val Leu
                                                      110
                                 105
Lys Glu Arg Arg Pro Leu Val Leu Cys Val Arg Glu Thr Pro Leu His
                             120
Leu Gly His Leu Arg Leu Met Thr Gln Ala Ala Glu Ile Gly Ala Val
Ile Met Pro Pro Val Pro Ala Phe Tyr His Arg Pro Gln Ser Leu Asp
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                     150
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165 170 175	
Thr Leu Pro Glu Asp Leu Phe Ala Arg Trp Gln Gly Ala 180 185	
<210> 155 <211> 25 <212> DNA <213> Artificial Sequence	
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